OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/061,201

DATE: 02/27/2002
TIME: 08:27:47

Input Set : D:\pto_PB0178.txt
Output Set: N:\CRF3\02272002\J061201.raw

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  3 <120> TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1
  5 <130> FILE REFERENCE: PB0178
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C--> 7 <141> CURRENT FILING DATE: 2002-01-30
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  8 <151> PRIOR FILING DATE: 2001-01-30
 10 <150> PRIOR APPLICATION NUMBER: PCT/US01/00667
 11 <151> PRIOR FILING DATE: 2001-01-30
 13 <150> PRIOR APPLICATION NUMBER: PCT/US01/00664
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 19 <150> PRIOR APPLICATION NUMBER: PCT/US01/00665
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 32 <151> PRIOR FILING DATE: 2001-05-23
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 35 <151> PRIOR FILING DATE: 2001-10-10
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 46 <400> SEQUENCE: 1
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 49 acaaggcaaa aattctgacg ttctcaagag accagctctg cccccgtggc tcaactgacc 180
 50 ctaccatgtg gacgctgctc ctccaggtgg gaactggagt tttgaaataa aatggatgat 240
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 53 cacaaagagc tgcggtgccc cgaatgcagg acgcctgtgt tttccaacat tgaggcgctg 420
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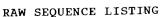
RAW SEQUENCE LISTING   DATE: 02/27/2002
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Output Set: **N:\CRF3\02272002\J061201.raw**

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60 cagctgcccc agccgccccc gctctgcagg gccctctaca acttcgacct acgaggcaag 840
61 gacaagagtg agaaccagga ttgcctgacc ttcctcaagg acgatatcat cactgtgatc 900
62 agccgagtgg atgagaactg ggcagaaggc aagttaggag ataaagtagg catcttccct 960
63 atcttgtttg tagagccaaa cctcaccgca agacaccttt tagagaagaa caaaggtcgc 1020
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86 cggagaaagt cagctcttgc caaggccaca accctggtgt ccactgcctc aggcacgcag 2400
87 accgtgtttc ccagcaaatg aacctacggg tgactttttcc tagaccccaa agaggtgaat 2460
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91 <211> LENGTH: 2190
92 <212> TYPE: DNA
93 <213> ORGANISM: Homo sapiens
95 <400> SEQUENCE: 2

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 97 gtcacagcca aagtcctccc ttgccagcac accttctgca aaccatgtct acagagggtt 120
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101 aaaagcagga ccaaccccag acgtctgcag gccagtcctt tccggctagt gcctaatgtc 360
102 agaatccaca tggatggggt gcctcgagca aaggccttat gcaactacag agggcagaat 420
103 cccggtgacc taaggtttaa taagggagat atcatccttc tccggagaca gcttgatgag 480
104 aattggtacc aggggggaaat caatggcatc agcgggaact tcccagccag ctccgtggaa 540
105 gtcatcaagc agctgcccca gccgcccccg ctctgcaggg ccctctacaa cttcgaccta 600
106 cgaggcaagg acaagagtga gaaccaggat tgcctgacct tcctcaagga cgatatcatc 660
107 actgtgatca gccgagtgga tgagaactgg gcagaaggca agttaggaga taaagtaggc 720
108 atcttccta tcttgtttgt agagccaaac ctcaccgcaa gacacctttt agagaagaac 780
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109 aaaggtcgcc agtcatcctg cacaaaaaac ctgtccctgg tgtcctcgtc ctccagaggc 840
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129 acctccggaa agcctgaaca gccagccacc ctcaaggcgt cccagcctga agcagcgtcc 2040
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131 acagacctcc ggagaaagtc agctcttgcc aaggccacaa ccctggtgtc cactgcctca 2160
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135 <211> LENGTH: 729
136 <212> TYPE: PRT
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139 <400> SEQUENCE: 3
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144             20                  25                  30
146 Cys Lys Pro Cys Leu Gln Arg Val Phe Lys Ala His Lys Glu Leu Arg
147         35                  40                  45
149 Cys Pro Glu Cys Arg Thr Pro Val Phe Ser Asn Ile Glu Ala Leu Pro
150     50                  55                  60
152 Ala Asn Leu Leu Leu Val Arg Leu Leu Asp Gly Val Arg Ser Gly Gln
153 65                  70                  75                  80
155 Ser Ser Gly Arg Gly Gly Ser Phe Arg Arg Pro Gly Thr Met Thr Leu
156                 85                  90                  95
158 Gln Asp Gly Arg Lys Ser Arg Thr Asn Pro Arg Arg Leu Gln Ala Ser
159             100                 105                 110
161 Pro Phe Arg Leu Val Pro Asn Val Arg Ile His Met Asp Gly Val Pro
162         115                 120                 125
164 Arg Ala Lys Ala Leu Cys Asn Tyr Arg Gly Gln Asn Pro Gly Asp Leu
165     130                 135                 140
167 Arg Phe Asn Lys Gly Asp Ile Ile Leu Leu Arg Arg Gln Leu Asp Glu
168 145                 150                 155                 160
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176 Arg Ala Leu Tyr Asn Phe Asp Leu Arg Gly Lys Asp Lys Ser Glu Asn
177             195                 200                 205
179 Gln Asp Cys Leu Thr Phe Leu Lys Asp Asp Ile Ile Thr Val Ile Ser
180         210                 215                 220
182 Arg Val Asp Glu Asn Trp Ala Glu Gly Lys Leu Gly Asp Lys Val Gly
183 225                 230                 235                 240
185 Ile Phe Pro Ile Leu Phe Val Glu Pro Asn Leu Thr Ala Arg His Leu
186                 245                 250                 255
188 Leu Glu Lys Asn Lys Gly Arg Gln Ser Ser Cys Thr Lys Asn Leu Ser
189             260                 265                 270
191 Leu Val Ser Ser Ser Ser Arg Gly Asn Thr Ser Thr Leu Arg Arg Gly
192             275                 280                 285
194 Pro Gly Ser Arg Arg Lys Val Pro Gly Gln Phe Ser Ile Thr Thr Ala
195     290                 295                 300
197 Leu Asn Thr Leu Asn Arg Met Val His Ser Pro Ser Gly Arg His Met
198 305                 310                 315                 320
200 Val Glu Ile Ser Thr Pro Val Leu Ile Ser Ser Ser Asn Pro Ser Val
201             325                 330                 335
203 Ile Thr Gln Pro Met Glu Lys Ala Asp Val Pro Ser Ser Cys Val Gly
204         340                 345                 350
206 Gln Val Ser Thr Tyr His Pro Ala Pro Val Ser Pro Gly His Ser Thr
207         355                 360                 365
209 Ala Val Val Ser Leu Pro Gly Ser Gln Gln His Leu Ser Ala Asn Met
210     370                 375                 380
212 Phe Val Ala Leu His Ser Tyr Ser Ala His Gly Pro Asp Glu Leu Asp
213 385                 390                 395                 400
215 Leu Gln Lys Gly Glu Gly Val Arg Val Leu Gly Lys Cys Gln Asp Gly
216             405                 410                 415
218 Trp Leu Arg Gly Val Ser Leu Val Thr Gly Arg Val Gly Ile Phe Pro
219             420                 425                 430
221 Asn Asn Tyr Val Ile Pro Ile Phe Arg Lys Thr Ser Ser Phe Pro Asp
222         435                 440                 445
224 Ser Arg Ser Pro Gly Leu Tyr Thr Thr Trp Thr Leu Ser Thr Ser Ser
225     450                 455                 460
227 Val Ser Ser Gln Gly Ser Ile Ser Glu Gly Asp Pro Arg Gln Ser Arg
228 465                 470                 475                 480
230 Pro Phe Lys Ser Val Phe Val Pro Thr Ala Ile Val Asn Pro Val Arg
231             485                 490                 495
233 Ser Thr Ala Gly Pro Gly Thr Leu Gly Gln Gly Ser Leu Arg Lys Gly
234         500                 505                 510
236 Arg Ser Ser Met Arg Lys Asn Gly Ser Leu Gln Arg Pro Leu Gln Ser
237         515                 520                 525
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240     530                 535                 540
242 Val Leu Arg Pro Gln Gln Phe Gln Phe Tyr Gln Pro Gln Gly Ile Pro
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251 Thr His Ser Ala Ala Ser Ser Leu Ile Met Glu Asp Lys Glu Ile Pro
252           595                   600                   605
254 Ile Lys Ser Glu Pro Leu Pro Lys Pro Pro Ala Ser Ala Pro Pro Ser
255       610                   615                   620
257 Ile Leu Val Lys Pro Glu Asn Ser Arg Asn Gly Ile Glu Lys Gln Val
258 625                   630                   635                   640
260 Lys Thr Val Arg Phe Gln Asn Tyr Ser Pro Pro Pro Thr Lys His Tyr
261               645                   650                   655
263 Thr Ser His Pro Thr Ser Gly Lys Pro Glu Gln Pro Ala Thr Leu Lys
264           660                   665                   670
266 Ala Ser Gln Pro Glu Ala Ala Ser Leu Gly Pro Glu Met Thr Val Leu
267       675                   680                   685
269 Phe Ala His Arg Ser Gly Cys His Ser Gly Gln Gln Thr Asp Leu Arg
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272 Arg Lys Ser Ala Leu Ala Lys Ala Thr Thr Leu Val Ser Thr Ala Ser
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286 acaaggcaaa aattctgacg ttctcaagag accagctctg cccccgtggc tcaactgacc 180
287 ctaccatgtg gacgctgctc ctccaggtgg gaactggagt tttgaaataa aatggatgat 240
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VERIFICATION SUMMARY
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DATE: 02/27/2002
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Output Set: **N:\CRF3\02272002\J061201.raw**

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L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date